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(54) Title: INVERSE LABELING METHOD FOR THE RAPID IDENTIFICATION OF MARKER/TARGET PROTEINS

erse Labeling Experimen

(57) Abstract: A novel procedure for performing protein labeling for comparative proteomics termed inverse labeling is provided for the profile of the profi for the rapid identification of marker or target proteins. With this method, to evaluate protein expression of a disease or a drug treated sample in comparison with a control sample, two converse collaborative labeling experiments are performed in parallel. In one experiment the perturbed sample (by disease or by drug treatment) is isotopically heavy-labeled, whereas, the control is isotopically heavy-labeled in the second experiment. Whem mixed and analyzed with its unlabeled or isotope light counterpart for differential comparison, a characteristic inverse labeling pattern is observed between the two parallel analyses for proteins that are differentially expressed to an apreciable level. In particularly useful embodiments, protein labeling is achieved through proteolytic ¹⁸O-incororation into peptides as a result of proteolysis performed in ¹⁸O-water, metabolic incoroporation of ¹⁵N(or ¹³C and ²H) into proteins, and chemically tagging proteins with an isotope-coded tag reagent such as an isotope-coded affinity tag reagent.

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INTERNATIONAL SEARCH REPORT



A. CLASSIFICATION OF SUBJECT MATTER IPC 7 G01N33/68

According to International Patent Classification (IPC) or to both national classification and IPC

Mnimum documentation searched (classification system followed by classification symbols) IPC 7-601N

Further documents are listed in the continuation of box C.

Documentation searched other than minimum documentation to the extent that such documents are included. In the fields searched

Electronic data base consulted during the international search (name of date base and, where practical, search terms used)

EPO-Internal, PAJ, WPI Data, MEDLINE, BIOSIS, EMBASE, SCISEARCH, CHEM ABS Data

Category * Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No
ROSE, K. ET AL: "A new mass-spectrometric C-terminal sequencing technique finds a similarity between gamma-interferon and alfa2-interferon and identifies a proteolytically clipped gamma-interferon that retains full antiviral activity" THE BIOCHEMICAL JOURNAL, vol. 215, no. 2, 1 November 1983 (1983-11-01), pages 273-277, XPOI079683 cited in the application the whole document	1-49

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Name and mailing address of the ISA European Patent ICHCe, P. B. 5818 Patentiaan 2 NL - 2280 HVF Ilijavijk Tel. (+31-70) 340-2040, Tx. 31 651 epc nl, Fax: (+31-70) 340-3016	Authorized officer Moreno de Vega, C

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X Patent family members are listed in annex.

INTERNATIONAL SEARCH REPORT

Interioral Application No PCT/EP 01/15228

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT Category * Citation of document, with indication, where appropriate, of the relevant passages Relevant to claim No. Α ODA, Y. ET AL: "Accurate quantitation of 1-49 protein expression and site-specific phosphorylation" PROC. NATL. ACAD. SCI., no. 96, 1999, pages 6591-6596, XP001080471 cited in the application the whole document JI, J. ET AL: "Strategy for qualitative 1-49 and quantitative analysis in proteomics based on signature peptides" JOURNAL OF CHROMATOGRAPHY. no. 745, 2000, pages 197-210, XP004215299 the whole document CHEN, X. ET AL: "Site-specific mass 1-49 tagging with stable isotopes in proteins for accurate and efficient protein identification" ANAL. CHEM., vol. 72, no. 6 15 March 2000 (2000-03-15), pages 1134-1143, XP002207456 cited in the application the whole document P,A WO 01 94935 A (MDS PROTEOMICS) 1-49 8 June 2001 (2001-06-08) claims 1-61 Ρ.Χ WANG, Y.K. ET AL: "Inverse 18 0 labeling 1-49 mass spectrometry for the rapid identification of marker/target proteins" ANALYTICAL CHEMISTRY vol. 73, no. 15, 1 August 2001 (2001-08-01), pages 3742-3750, XP001061446 USA the whole document

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Patent document cited in search report		Publication date		Patent family member(s)		Publication date	
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